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Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: [year=2008; month=7; day=31; hr=16; min=30; sec=47; ms=911;]

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Application No: 08822963 Version No: 1.0

Input Set:

Output Set:

Started: 2008-07-31 15:09:05.460
Finished: 2008-07-31 15:09:06.199
Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 739 ms
Total Warnings: 7
Total Errors: 0
No. of SeqIDs Defined: 16
Actual SeqID Count: 16

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (2)
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W 402	Undefined organism found in <213> in SEQ ID (9)
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W 402	Undefined organism found in <213> in SEQ ID (14)

<110> Liu, Dakai

Rabbani, Elazar

<120> VECTORS AND VIRAL VECTORS, AND PACKAGING CELL LINES FOR
PROPOGATING SAME

<130> ENZ-56SequenceListing.110398

<140> 08822963

<141> 2008-07-31

<160> 16

<170> PatentIn Ver. 2.0

<210> 1

<211> 9

<212> DNA

<213> Bacteriophage lambda

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 1

tatcaccgc

9

<210> 2

<211> 9

<212> DNA

<213> bacteriophage 434

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 2

acaagaaaa

9

<210> 3

<211> 10

<212> DNA

<213> Escherichia coli

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 3

gtactagtta

10

<210> 4

<211> 8

<212> DNA

<213> Escherichia coli

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 4

agacgtct

8

<210> 5

<211> 24

<212> DNA

<213> Escherichia coli

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 5

tggaattgtg agcggataac aatt

24

<210> 6

<211> 4

<212> DNA

<213> Drosophila melanogaster

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 6

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4

<210> 7

<211> 9

<212> DNA

<213> MAT alpha 2 yeast

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 7

catgtaatt

9

<210> 8

<211> 13

<212> DNA

<213> Escherichia coli

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 8

aaaagtgtga cat

13

<210> 9

<211> 11

<212> DNA

<213> GAL4 yeast

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 9

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11

<210> 10

<211> 12

<212> DNA

<213> Papillomavirus sylvilagi

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

<400> 10

accgacgtcg gt

12

<210> 11

<211> 6

<212> DNA

<213> GCN4 yeast

<220>

<223> Description of Artificial Sequence:nucleic acid,

double stranded, linear topology

<400> 11

atgatac

6

<210> 12

<211> 9

<212> DNA

<213> zif268 murine

<220>

<223> Description of Artificial Sequence:nucleic acid,

double stranded, linear topology

<400> 12

gcgtgggacg

9

<210> 13

<211> 9

<212> DNA

<213> human glucocorticoid

<220>

<223> Description of Artificial Sequence:nucleic acid,

double stranded, linear topology

<400> 13

cagaacatc

9

<210> 14

<211> 8

<212> DNA

<213> tfiid

<220>

<223> Description of Artificial Sequence:nucleic acid,
double stranded, linear topology

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tatataaa

8

<210> 15

<211> 319

<212> DNA

<213> murine leukemia virus

<220>

<223> Description of Artificial Sequence:nucleic acid,
single stranded, linear topology

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taagcagttc ctgccccggc tcagggccaa gaacagatgg tccccagatg cgggccagcc 180

ctcagcagtt tctagagaac catcagatgt ttccagggtg cccaaggac ctgaaatgac 240

cctgtgcctt atttgaacta accaatcagt tcgcttctcg cttctgttcg cgcgcttctg 300

ctccccgagc tcaataaaa

319

<210> 16

<211> 319

<212> DNA

<213> murine leukemia virus

<220>

<223> Description of Artificial Sequence:nucleic acid,

single stranded, linear topology

<400> 16

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ctcgcgccag ccgaactgtt cgccaggctc aaggcgcgca tgcccgaacg cgaggatctc 180

gtcgtgactt tctagagaac catcagatgt ttccaggggtg cccaaggac ctgaaatgac 240

cctgtgcctt atttgaacta accggtcagt tcgcttctcg cttctgttcg cgcgcttctg 300

ctccccgagc tcagctgcg 319